#8

SEQUENCE LISTING

<110> Watowich, Stanley J.
Weaver, Scott C.
Davey, Robert A.

<120> Drug Discovery Methods

<130> 265.00260101

<140> US 09/981,286
<141> 2001-10-15

<150> US 60/240,187
<151> 2000-10-13

<160> 36

<170> PatentIn version 3.0

<210> 1
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<213> VENEZUELAN EQUINE ENCEPHALITIS VIRUS

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Val Met Lys Leu Glu Ser Asp Lys Thr Phe Pro Ile Met Leu Glu Gly
1 5 10 15

Lys Ile Asn Gly Tyr Ala Cys Val Val Gly Gly Lys Leu Phe Arg Pro
20 25 30

Met His Val Glu Gly Lys Ile Asp Asn Asp Val Leu Ala Ala Leu Lys
35 40 45

Thr Lys Lys Ala Ser Lys Tyr Asp Leu Glu Tyr Ala Asp Val Pro Gln
50 55 60

Asn Met Arg Ala Asp Thr Phe Lys Tyr Thr His Glu Lys Pro Gln Gly
65 70 75 80

Tyr Tyr Ser Trp His His Gly Ala Val Gln Tyr Glu Asn Gly Arg Phe
85 90 95

Thr Val Pro Lys Gly Val Gly Ala Lys Gly Asp Ser Gly Arg Pro Ile
100 105 110

Leu Asp Asn Gln Gly Arg Val Val Ala Ile Val Leu Gly Gly Val Asn
115 120 125

Glu Gly Ser Arg Thr Ala Leu Ser Val Val Met Trp Asn Glu Lys Gly
130 135 140

Val Thr Val Lys Tyr Thr Pro Glu Asn Cys Glu Gln Trp
145 150 155

<210> 2

<211> 11

<212> PRT

<213> ARTIFICIAL

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<223> Cell-permeant polypeptide

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Tyr Gly Arg Lys Lys Arg Arg Gln Arg Arg Arg
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<210> 3

<211> 16

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<213> ARTIFICIAL

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Arg Gln Ile Lys Ile Trp Phe Gln Asn Arg Arg Met Lys Trp Lys Lys
1 5 10 15

<210> 4

<211> 16

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<223> Cell-permeant polypeptide

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Arg Gln Ile Lys Ile Trp Phe Pro Asn Arg Arg Met Lys Trp Lys Lys
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<211> 16

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<223> Cell-permeant polypeptide

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Arg Gln Pro Lys Ile Trp Phe Pro Asn Arg Arg Pro Lys Trp Lys Lys
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<211> 525

<212> DNA

<213> ARTIFICIAL

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<223> Nucleotide sequence encoding tat-CCD

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acgttcccaa tcatgttgga agggaagata aacggctacg cttgtgtggt cggagggaag 120

ttattcaggc cgatgcatgt ggaaggcaag atcgacaacg acgttctggc cgcgcttaag 180

acgaagaaag catccaaata cgatcttgag tatgcagatg tgccacagaa catgcgggcc 240

gatacattca aatacaccca tgagaaaccc caaggctatt acagctggca tcatggagca 300

gtccaatatg aaaatgggcg tttcacggtg ccgaaaggag ttggggccaa gggagacagc 360

ggacgaccca ttctggataa ccagggacgg gtggtcgcta ttgtgctggg aggtgtgaat 420

gaaggatcta ggacagccct ttcagtcgtc atgtggaaca agcttggatc ttctctcgag 480

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<223> Amino acid sequence of tat-CCD

<400> 7

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Glu Ser Asp Lys Thr Phe Pro Ile Met Leu Glu Gly Lys Ile Asn Gly
            20                  25                  30

Tyr Ala Cys Val Val Gly Gly Lys Leu Phe Arg Pro Met His Val Glu
        35                  40                  45

Gly Lys Ile Asp Asn Asp Val Leu Ala Ala Leu Lys Thr Lys Lys Ala
    50                  55                  60

Ser Lys Tyr Asp Leu Glu Tyr Ala Asp Val Pro Gln Asn Met Arg Ala
65                  70                  75                  80

Asp Thr Phe Lys Tyr Thr His Glu Lys Pro Gln Gly Tyr Tyr Ser Trp
                85                  90                  95

His His Gly Ala Val Gln Tyr Glu Asn Gly Arg Phe Thr Val Pro Lys
            100                 105                 110

Gly Val Gly Ala Lys Gly Asp Ser Gly Arg Pro Ile Leu Asp Asn Gln
        115                 120                 125

Gly Arg Val Val Ala Ile Val Leu Gly Gly Val Asn Glu Gly Ser Arg
    130                 135                 140

Thr Ala Leu Ser Val Val Met Trp Asn Glu Lys Gly Val Thr Val Lys
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Tyr Thr Pro Glu Asn Cys Glu Gln Trp
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<210> 8

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<212> PRT

<213> BOS TAURUS

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Lys Glu Thr Ala Ala Ala Lys Phe Glu Arg Gln His Met Asp Ser Ser
1               5                  10                  15
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Thr Ser Ala Ala Ser Ser Ser Asn Tyr Cys Asn Gln Met Met Lys Ser
20 25 30

Arg Asn Leu Thr Lys Asp Arg Cys Lys Pro Val Asn Thr Phe Val His
35 40 45

Glu Ser Leu Ala Asp Val Gln Ala Val Cys Ser Gln Lys Asn Val Ala
50 55 60

Cys Lys Asn Gly Gln Thr Asn Cys Tyr Gln Ser Tyr Ser Thr Met Ser
65 70 75 80

Ile Thr Asp Cys Arg Glu Thr Gly Ser Ser Lys Tyr Pro Asn Cys Ala
85 90 95

Tyr Lys Thr Thr Gln Ala Asn Lys His Ile Ile Val Ala Cys Glu Gly
100 105 110

Asn Pro Tyr Val Pro Val His Phe Ala Ala Ser Val
115 120

<210> 9

<211> 37

<212> DNA

<213> ARTIFICIAL

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<223> Primer

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<210> 10

<211> 42

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<213> ARTIFICIAL

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<223> Any amino acid

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<213> VENEZUELAN EQUINE ENCEPHALITIS VIRUS

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aacgacgttc tggccgcgct taagacgaag aaagcatcca aatacgatct tgagtatgca 180

gatgtgccac agaacatgcg ggccgataca ttcaaataca cccatgagaa accccaaggc 240

tattacagct ggcatcatgg agcagtccaa tatgaaaatg ggcgtttcac ggtgccgaaa 300

ggagttgggg ccaagggaga cagcggacga cccattctgg ataaccaggg acgggtggtc 360

gctattgtgc tgggaggtgt gaatgaagga tctaggacag ccctttcagt cgtcatgtgg 420

aacgagaagg gagttaccgt gaagtatact ccggagaact gcgagcaatg gtaatga 477

<210> 13

<211> 43

<212> DNA

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agctaggaat tcggatccca tatgtacggt cgtaaaaaac gtc 43

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ctagctaagc ttgttccaca tgacgactga aag 33

<210> 15

<211> 36

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ctagctgcgg ccgctcatta ccattgctcg cagttc 36

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<210> 17

<211> 50

<212> DNA

<213> ARTIFICIAL

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<211> 51

<212> DNA

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<222> (15)..(16)

<223> A, T, G, or C

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agctttccgg tggtnnkggt ggttccc 27

<210> 20

<211> 10

<212> DNA

<213> ARTIFICIAL

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<223> Primer

<400> 20
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<210> 21

<211> 14

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<400> 21
tcgagggaac cacc 14

<210> 22

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<400> 22
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<222> (15)..(16)

<223> A, T, G, or C

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<211> 19

<212> DNA

<213> ARTIFICIAL

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<223> Primer

<400> 27
cgatctcgag ggaaccatc 19

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<211> 57

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<222> (22)..(23)

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<223> A, T, G, or C

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<223> A, T, G, or C

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<222> (31)..(32)

<223> A, T, G, or C

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<222> (34)..(35)

<223> A, T, G, or C

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<211> 21

<212> DNA

<213> ARTIFICIAL

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<400> 29
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<210> 30

<211> 18

<212> DNA

<213> ARTIFICIAL

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<223> A, T, G, or C

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<222> (4)..(5)

<223> A, T, G, or C

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<221> misc_feature

<222> (7)..(8)

<223> A, T, G, or C

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<222> (10)..(11)

<223> A, T, G, or C

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<222> (13)..(14)

<223> A, T, G, or C

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<222> (16)..(17)

<223> A, T, G, or C

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<212> DNA

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<223> Adaptein-1 nucleotide sequence

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tacgcttgtg tggtcggagg gaagttattc aggccgatgc atgtggaagg caagatcgac 120

aacgacgttc tggccgcgct taagacgaag aaagcatcca aatacgatct tgagtatgca 180

gatgtgccac agaacatgcg ggccgataca ttcaaataca cccatgagaa accccaaggc 240

tattacagct ggcatcatgg agcagtccaa tatgaaaatg ggcgtttcac ggtgccgaaa 300

ggagttgggg ccaagggaga cagcggacga cccattctgg ataaccaggg acgggtggtc 360

gctattgtgc tgggaggtgt gaatgaagga tctaggacag ccctttcagt cgtcatgtgg 420

aacaagcttt ctccacatta tgctcaactc gagggagtta ccgtgaagta tactccggag 480

aactgcgagc aatggtaatg agc 503

<210> 32

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<212> DNA

<213> ARTIFICIAL

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<223> Adaptein-2 nucleotide sequence

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aacgacgttc tggccgcgct taagacgaag aaagcatcca aatacgatct tgagtatgca 180

gatgtgccac agaacatgcg ggccgataca ttcaaataca cccatgagaa accccaaggc 240

tattacagct ggcatcatgg agcagtccaa tatgaaaatg ggcgtttcac ggtgccgaaa 300

ggagttgggg ccaagggaga cagcggacga cccattctgg ataaccaggg acgggtggtc 360

gctattgtgc tgggaggtgt gaatgaagga tctaggacag ccctttcagt cgtcatgtgg 420

aacaagctta gaagcggtac tcaatggctc gagggagtta ccgtgaagta tactccggag 480

aactgcgagc aatggtaatg agc 503

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<212> PRT

<213> ARTIFICIAL

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<223> Adaptein-1 amino acid sequence

<400> 33

Val Met Lys Leu Glu Ser Asp Lys Thr Phe Pro Ile Met Leu Glu Gly
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Lys Ile Asn Gly Tyr Ala Cys Val Val Gly Gly Lys Leu Phe Arg Pro
20 25 30

Met His Val Glu Gly Lys Ile Asp Asn Asp Val Leu Ala Ala Leu Lys
35 40 45

Thr Lys Lys Ala Ser Lys Tyr Asp Leu Glu Tyr Ala Asp Val Pro Gln
50 55 60

Asn Met Arg Ala Asp Thr Phe Lys Tyr Thr His Glu Lys Pro Gln Gly
65 70 75 80

Tyr Tyr Ser Trp His His Gly Ala Val Gln Tyr Glu Asn Gly Arg Phe
85 90 95

Thr Val Pro Lys Gly Val Gly Ala Lys Gly Asp Ser Gly Arg Pro Ile
100 105 110

Leu Asp Asn Gln Gly Arg Val Val Ala Ile Val Leu Gly Gly Val Asn

115 120 125

Glu Gly Ser Arg Thr Ala Leu Ser Val Val Met Trp Asn Lys Leu Ser
130 135 140

Pro His Tyr Ala Gln Leu Glu Gly Val Thr Val Lys Tyr Thr Pro Glu
145 150 155 160

Asn Cys Glu Gln Trp
165

<210> 34

<211> 165

<212> PRT

<213> ARTIFICIAL

<220>

<223> Adaptein-2 amino acid sequence

<400> 34

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Lys Ile Asn Gly Tyr Ala Cys Val Val Gly Gly Lys Leu Phe Arg Pro
20 25 30

Met His Val Glu Gly Lys Ile Asp Asn Asp Val Leu Ala Ala Leu Lys
35 40 45

Thr Lys Lys Ala Ser Lys Tyr Asp Leu Glu Tyr Ala Asp Val Pro Gln
50 55 60

Asn Met Arg Ala Asp Thr Phe Lys Tyr Thr His Glu Lys Pro Gln Gly
65 70 75 80

Tyr Tyr Ser Trp His His Gly Ala Val Gln Tyr Glu Asn Gly Arg Phe
85 90 95

Thr Val Pro Lys Gly Val Gly Ala Lys Gly Asp Ser Gly Arg Pro Ile
100 105 110

Leu Asp Asn Gln Gly Arg Val Val Ala Ile Val Leu Gly Gly Val Asn
115 120 125

Glu Gly Ser Arg Thr Ala Leu Ser Val Val Met Trp Asn Lys Leu Arg
130 135 140

Ser Gly Thr Gln Trp Leu Glu Gly Val Thr Val Lys Tyr Thr Pro Glu
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Asn Cys Glu Gln Trp
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<211> 58

<212> DNA

<213> ARTIFICIAL

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<211> 42

<212> DNA

<213> ARTIFICIAL

<220>

<223> Primer

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